

# INKA, an integrative data analysis pipeline for phosphoproteomic inference of active kinases

Robin Beekhof, Carolien van Alphen, Alex A Henneman, Jaco C Knol, Thang V Pham, Frank Rolfs, Mariette Labots, Evan Henneberry, Tessa YS Le Large, Richard R de Haas, Sander R Piersma, Valentina Vurchio, Andrea Bertotti, Livio Trusolino, Henk MW Verheul & Connie R Jimenez

**Correction to:** *Mol Syst Biol* (2019) 15: e8250. DOI 10.15252/msb.20188250 | Published online 12 April 2019

The authors recently noticed that the affiliations for Valentina Vurchio, Andrea Bertotti, and Livio Trusolino were listed incorrectly. The correct author affiliation is shown here and has been corrected in the original article. The authors apologize for this error.

Valentina Vurchio<sup>3</sup>, Andrea Bertotti<sup>3,4</sup>, Livio Trusolino<sup>3,4</sup>

<sup>3</sup>Candiolo Cancer Institute - FPO IRCCS, Candiolo, Torino, Italy

<sup>4</sup>Department of Oncology, University of Torino, Candiolo, Torino, Italy